Appendix I: Alignment of SEO ID NO: 7 with the chloramphenical resistance gene of Borges et al.

```
<!--StartFragment-->RESULT 3
AAS00462
ID
    AAS00462 standard; DNA; 2003 BP.
ХX
AC
    AAS00462;
XX
DΤ
    11-JUN-2007 (revised)
DT
    16-MAY-2001 (first entry)
XX
DE
    Plasmid pLOI2225 useful for chromosomal integration of heterologous DNA.
XX
KW
     Plasmid; vector; antibiotic resistance; ethanol; alcohol dehydrogenase;
KW
     adhB; pyruvate decarboxylase; pdc; chloramphenicol acetyl transferase;
KW
     cat; regulatory element; adhE; chromosomal integration; circular; cyclic;
KW
     pLOI2225; pLOI2222; pSG76-C; chloramphenicol resistance;
KW
    FRT recombining site; ds.
XX
os
     Synthetic.
XX
FH
     Kev
                     Location/Qualifiers
FT
     CDS
                     complement (1045, .1704)
FT
                     /*tag= a
FT
                     /note= "Chloramphenicol-resistance gene"
XX
PN
     WO200118222-A1.
XX
PD
     15-MAR-2001.
XX
PF
     18-AUG-2000; 2000WO-US022700.
XX
PR
     07-SEP-1999:
                    99US-00390479.
XX
PA
    (UYFL ) UNIV FLORIDA.
XX
PΙ
     Borges AC, Zaldivar J, Morales FM, Jimenez AM, Ingram LO;
XX
DR
    WPI: 2001-235205/24.
DR
     PC:NCBI; qi6467484.
XX
PT
    Novel nucleic acid construct for integrating heterologous nucleic acid
PT
    sequences into genome or chromosome of host cells, has passenger and
PT
    marker sequences, in which marker sequence is flanked by recombining
PT
    sites.
XX
PS
    Claim 28; Page 59-60; 85pp; English.
XX
CC
    The present sequence for plasmid pLOI2225 which is constructed from the
CC
    plasmids pLOI2222 and pSG76-C is 1 of 7 novel plasmid constructs
CC
    (AAS00460-AAS00466) comprising a marker sequence such as an antibiotic
CC
     resistance gene, in which the marker sequence is flanked by two FRT
CC
     recombining sites. One of these plasmids (pLOI02231) also comprises a
CC
    passenger sequence. The passenger sequence can include an ethanologenic
CC
    gene such as alcohol dehydrogenase (preferably adhB) or pyruvate
CC
    decarboxylase (pdc), another gene such as chloramphenicol acetyl
CC
    transferase (cat), a regulatory element such as a promoter or IRES
CC
    (internal ribosomal entry site) or a quide sequence such as adhE. All the
CC
    plasmids are useful for integrating a nucleic acid construct into the
CC
    genome of a cell. Plasmid pLOI02231 is useful for producing ethanol by,
CC
    transforming an ethanologenic cell with the plasmid and contacting the
CC
     cell with a substrate which can be fermented into ethanol, where
CC
     expression of the passenger sequence results in the production of
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ethanol. The recombinant ethanologenic host transformed with the plasmid
CC
    has improved properties including increased ability to produce ethanol,
    depolymerisation for a particular substrate and increased tolerance to a
CC
CC
    higher level of ethanol
CC
CC
    Revised record issued on 11-JUN-2007 : Enhanced with precomputed
CC
    information from BOND.
XX
so
    Sequence 2003 BP; 558 A; 425 C; 436 G; 584 T; 0 U; 0 Other;
 Query Match
                       100.0%; Score 1069; DB 1; Length 2003;
 Best Local Similarity
                     100.0%; Pred. No. 0;
 Matches 1069; Conservative
                             0; Mismatches
                                              0; Indels
                                                          0; Gaps
Qу
          1 GCAAAAATTAAAAATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAAACTTGGTCTG 60
         935 GCAAAAATTAAAAATGAAGTTTTAAATCAATCTAAAGTATATATGAGTAAACTTGGTCTG 994
Dh
         61 ACAGTTACCAATGCTTAATCAGTGAGGCACCAATAACTGCCTTAAAAAAATTACGCCCCG 120
Qy
         Db
        121 CCCTGCCACTCATCGCAGTACTGTTGTAATTCATTAAGCATTCTGCCGACATGGAAGCCA 180
QУ
Dh
        1055 CCCTGCCACTCATCGCAGTACTGTTGTAATTCATTAAGCATTCTGCCGACATGGAAGCCA 1114
        181 TCACAGACGCATGATGAACCTGAATCGCCAGCGCATCAGCACCTTGTCGCCTTGCGTA 240
Qy
            1115 TCACAGACGGCATGATGAACCTGAATCGCCAGCGGCATCAGCACCTTGTCGCCTTGCGTA 1174
Dh
         241 TAATATTTGCCCATGGTGAAAACGGGGGGCGAAGAAGTTGTCCATATTGGCCACGTTTAAA 300
Ov
             Db
        1175 TAATATTTGCCCATGGTGAAAACGGGGGCGAAGAAGTTGTCCATATTGGCCACGTTTAAA 1234
         301 TCAAAACTGGTGAAACTCACCCAGGGATTGGCTGAGACGAAAAACATATTCTCAATAAAC 360
Qу
Db
        1235 TCAAAACTGGTGAAACTCACCCAGGGATTGGCTGAGACGAAAAACATATTCTCAATAAAC 1294
        361 CCTTTAGGGAAATAGGCCAGGTTTTCACCGTAACACGCCACATCTTGCGAATATATGTGT 420
QУ
Dh
        1295 CCTTTAGGGAAATAGGCCAGGTTTTCACCGTAACACGCCACATCTTGCGAATATATGTGT 1354
Οv
         421 AGAAACTGCCGGAAATCGTCGTGGTATTCACTCCAGAGCGATGAAAACGTTTCAGTTTGC 480
        1355 AGAAACTGCCGGAAATCGTCGTGGTATTCACTCCAGAGCGATGAAAACGTTTCAGTTTGC 1414
        481 TCATGGAAAACGGTGTAACAAGGGTGAACACTATCCCATATCACCAGCTCACCGTCTTTC 540
Qу
        1415 TCATGGAAAACGGTGTAACAAGGGTGAACACTATCCCATATCACCAGCTCACCGTCTTTC 1474
Db
         541 ATTGCCATACGGATTTCGGATGAGCATTCATCAGGCGGGCAAGAATGTGAATAAAGGCC 600
QУ
        1475 ATTGCCATACGGAATTTCGGATGAGCATTCATCAGGCGGGCAAGAATGTGAATAAAGGCC 1534
Db
        601 GGATAAAACTTGTGCTTATTTTTCTTTACGGTCTTTAAAAAGGCCGTAATATCCAGCTGA 660
             1535 GGATAAAACTTGTGCTTATTTTTCTTTACGGTCTTTAAAAAGGCCGTAATATCCAGCTGA 1594
        661 ACGGTCTGGTTATAGGTACATTGAGCAACTGACTGAAATGCCTCAAAATGTTCTTTACGA 720
        1595 ACGGTCTGGTTATAGGTACATTGAGCAACTGACTGAAATGCCTCAAAATGTTCTTTACGA 1654
Db
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Qy	721	TGCCATTGGGATATATCAACGGTGGTATATCCAGTGATTTTTTTCTCCATTTTAGCTTCC	780
Db	1655	${\tt TGCCATTGGGATATATCAACGGTGGTATATCCAGTGATTTTTTTCTCCATTTTAGCTTCC}$	1714
QУ	781	TTAGCTCCTGAAAATCTCGATAACTCAAAAAATACGCCCGGTAGTGATCTTATTTCATTA	840
Db	1715		1774
Qy	841	TGGTGAAAGTTGGAACCTCTTACGTGCCGATCAACGTCTCATTTTCGCCAAAAGTTGGCC	900
Db	1775	${\tt TGGTGAAAGTTGGAACCTCTTACGTGCCGATCAACGTCTCATTTTCGCCAAAAGTTGGCC}$	1834
Qy	901	CAGGGCTTCCCGGTATCAACAGGGACCCAGGATTTATTTA	960
Db	1835	${\tt CAGGGCTTCCCGGTATCAACAGGGACACCAGGATTTATTT$	1894
Qy	961	GTCACAGGTATTTATTCGGCGCAAAGTGCGTCGGGTGATGCTGCCAACTTACTGATTTAG	1020
Db	1895		1954
Qy	1021	TGTATGATGGTGTTTTTGAGGTGCTCCAGTGGCTTCTGTTTCTATCAGC 1069	
Db	1955		

<!--EndFragment-->